

GenCore version 5.1.3  
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OM nucleic - nucleic search, using sw model

Run on: December 13, 2002, 15:41:52 : Search time 268 seconds  
(without alignments)  
1378.089 Million cell updates/sec

Title: US-09-659-737A-1

Perfect score: 164

Sequence: 1 gcaccgggacatcaaggag.....tgctggtggtgccccagaag 164

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 1125999159 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

N.Geneseq\_101002.\*  
1: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1980.DAT.\*  
2: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1981.DAT.\*  
3: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1982.DAT.\*  
4: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1983.DAT.\*  
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8: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA1987.DAT.\*  
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21: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2000.DAT.\*  
22: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001A.DAT.\*  
23: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2001B.DAT.\*  
24: /SIDS2/gcgdata/geneseq/geneseq-emb1/NA2002.DAT.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

| Result No. | Query Match % | Score | Length | ID | Description                  |
|------------|---------------|-------|--------|----|------------------------------|
| 1          | 164           | 100.0 | 164    | 22 | AAF30487 Human MLK4 partial  |
| 2          | 152.8         | 93.2  | 2157   | 22 | AAH46913 CDNA encoding huma  |
| 3          | 152.8         | 93.2  | 3066   | 24 | ABQ86165 Novel human gene.   |
| 4          | 152.8         | 93.2  | 3111   | 24 | ABN86357 Novel human protei  |
| 5          | 152.8         | 93.2  | 3518   | 22 | ABN86358 Novel human protei  |
| 6          | 94.2          | 57.4  | 3141   | 22 | AAAD18824 Human kinase (PKIN |
| 7          | 94.2          | 57.4  | 3538   | 24 | AAAD34309 Human PKIN-12 cDNA |
| 8          | 87            | 53.0  | 444    | 22 | AAI85458 Human polynucleoti  |
| 9          | 83            | 50.6  | 3558   | 24 | ABK83874 Human cDNA differe  |

|    |      |      |        |    |                               |
|----|------|------|--------|----|-------------------------------|
| 10 | 83   | 50.6 | 3558   | 24 | AAH36139 Human mitogen acti   |
| 11 | 73.4 | 44.8 | 3454   | 24 | ABL70018 Pancreas cancer re   |
| 12 | 72.4 | 44.1 | 7084   | 22 | ABA17177 Human nervous syst   |
| 13 | 72.4 | 44.1 | 7084   | 22 | AAK73916 Human immune/haema   |
| 14 | 72   | 43.9 | 404    | 14 | AAO49755 pTK gene LptK4 par   |
| 15 | 72   | 43.9 | 404    | 16 | AAO49755 Protein tyrosine-k   |
| 16 | 62   | 37.8 | 137    | 14 | AAO49751 pTK gene LptK4 par   |
| 17 | 62   | 37.8 | 137    | 16 | AAO49751 Protein tyrosine-k   |
| 18 | 57.4 | 35.0 | 3072   | 23 | ABL04365 Drosophila melanog   |
| 19 | 57.4 | 35.0 | 6721   | 23 | ABL04365 Drosophila melanog   |
| 20 | 33.2 | 20.2 | 3365   | 24 | ABR84403 Human cDNA differe   |
| 21 | 33.2 | 20.2 | 3389   | 16 | AAAT01031 Human leucine-zipp  |
| 22 | 33.2 | 20.2 | 3389   | 18 | AAAT89349 Human leucine-zipp  |
| 23 | 33   | 20.1 | 169    | 21 | AAA45132 Mouse secreted exp   |
| 24 | 32.2 | 19.6 | 3427   | 24 | AB199250 Mouse ischaemic co   |
| 25 | 32   | 19.5 | 1008   | 20 | AAAX07105 Staphylococcus aur  |
| 26 | 32   | 19.5 | 1008   | 23 | AAAS4418 Staphylococcus aur   |
| 27 | 32   | 19.5 | 3647   | 18 | AAV74332 Staphylococcus aur   |
| 28 | 31.4 | 19.1 | 604    | 22 | AAAS5634 Human cardiovascular |
| 29 | 31.4 | 19.1 | 7116   | 23 | ABL03328 Drosophila melanog   |
| 30 | 31   | 18.9 | 751    | 14 | AAQ43394 Drosophila melanog   |
| 31 | 31   | 18.9 | 3109   | 23 | ABL29755 Drosophila melanog   |
| 32 | 31   | 18.9 | 12049  | 23 | ABL29754 Human cDNA differe   |
| 33 | 30.8 | 18.7 | 2161   | 22 | ABR84377 Human secreted p     |
| 34 | 30.6 | 18.7 | 5073   | 21 | AAA09254 Human alpha-2-delt   |
| 35 | 30.6 | 18.7 | 5073   | 22 | AAA09254 Human cardiovascular |
| 36 | 30.6 | 18.7 | 5557   | 22 | AAAS3573 Human cardiovascular |
| 37 | 30.6 | 18.7 | 5557   | 22 | AAAS3573 Human immune/haema   |
| 38 | 30.6 | 18.7 | 5557   | 22 | AAK73043 Human alpha-2-delt   |
| 39 | 30.6 | 18.7 | 5713   | 22 | AAK09278 Human oestrogen re   |
| 40 | 30.6 | 18.7 | 465237 | 24 | ABO87691 Human oestrogen re   |
| 41 | 30.6 | 18.7 | 465237 | 24 | ABR90193 Human cDNA differe   |
| 42 | 30.4 | 18.5 | 1540   | 24 | ABR63582 cDNA sequence enco   |
| 43 | 30.4 | 18.5 | 1597   | 21 | AAK46156 Genomic sequence #   |
| 44 | 30.4 | 18.5 | 10378  | 23 | ABK43051 Human cDNA sequenc   |
| 45 | 30.2 | 18.4 | 1542   | 22 | AAH16949                      |

#### ALIGNMENTS

RESULT 1  
AAF30487  
ID AAF30487 standard; cDNA; 164 BP.

AC AAF30487;  
DT 29-MAY-2001 (first entry)  
DE Human MLK4 partial cDNA.

KW MLK4; human; c-Jun N-terminal kinase kinase kinase; JNKKK;  
KW protein kinase; ultraviolet radiation; skin damage; inflammation;  
KW psoriasis; radioprotective; antiinflammatory; antiproliferative;  
KW vunerary; ss.

OS Homo sapiens.

Key Location/Qualifiers  
CDS 2..163  
FT /\*tag= a  
FT /partial

FN EF1085093-A2.

PD 21-MAR-2001.

FF 12-SEP-2000; 2000EP-0307866.

PR 20-SEP-1999; 990S-0155029.

PA (UUNY ) UNIV NEW YORK STATE.



DT 10-SEP-2002 (first entry)  
 XX Novel human gene. SEQ ID 36.  
 DE Human; cytostatic; vulnerary; antiarteriosclerotic; antiparkinsonian;  
 XX neurotropic; neuroprotective; immunosuppressive; haemostatic;  
 KW antiinflammatory; cardiant; antitumor; virucide; antithyroid;  
 KW cerebroprotective; anorectic; metabolic; vaccine; cancer; infection;  
 KW wound healing disease; atherosclerosis; Parkinson's disease;  
 KW Alzheimer's disease; autoimmune disorder; haematopoietic disorder;  
 KW inflammation; neoplastic disease; nervous system disorder;  
 KW cardiovascular disorders; pancreatitis; respiratory disorder;  
 KW hyperproliferation; systemic autoimmune disease; hyper-immunity;  
 KW developmental abnormality; gastrointestinal ulceration; neuropathy;  
 KW haematological abnormality; metabolic disease; sperm dysfunction;  
 KW thyroid disorder; hypothyroidism; brain damage; colitis;  
 KW cone photo transduction deficiency; neurological disease; stroke;  
 KW angio genesis; ovulation disorder; spinal cord; thyroid gland; heart;  
 KW trachea; thymus; lymph node; muscular system; obesity; anorexia;  
 KW growth abnormality; precocious puberty; gene; ss.  
 XX Homo sapiens.  
 OS WO200250105-A1.  
 XX 27-JUN-2002.  
 XX 17-DEC-2001; 2001WO-US49232.  
 XX 19-DEC-2000; 2000US-256710P.  
 PR 20-DEC-2000; 2000US-257048P.  
 PR 09-JAN-2001; 2001US-260482P.  
 PR 30-JAN-2001; 2001US-264922P.  
 PR 06-FEB-2001; 2001US-266797P.  
 PR 19-MAR-2001; 2001US-276988P.  
 PR 04-APR-2001; 2001US-281535P.  
 PR 08-MAY-2001; 2001US-289622P.  
 XX (SMIK ) SMITHKLINE BEECHAM CORP.  
 PA (SMIK ) SMITHKLINE BEECHAM PLC.  
 PA (GLAX ) GLAXO GROUP LTD.  
 XX Agarwal P, Birkeland M, Cogswell JP, Kabnick KF, Lai Y;  
 PI Martensen SA, Rizvi SK, Smith RF, Strum JC, Xie Q;  
 XX WPI: 2002-508784/54.  
 DR P-PSDB; ABP61000.  
 XX Secreted proteins and polynucleotides useful as vaccines for preventing  
 PI or treating various diseases e.g. cancer, wounds, atherosclerosis,  
 PT Parkinson's disease, Alzheimer's disease, infection, autoimmune  
 PT disorder -  
 XX Claim 2(a): Page 249; 335pp; English.  
 XX The invention relates to an isolated polypeptide with signal sequences  
 CC which allow it to be secreted extracellularly or membrane associated.  
 CC The activity of polypeptides of the invention may be described as,  
 CC cytostatic, vulnerary, antiarteriosclerotic, antiparkinsonian, neurotropic,  
 CC neuroprotective, immunosuppressive, haemostatic, antiinflammatory,  
 CC cardiant, antitumor, virucide, antithyroid, cerebroprotective, anorectic,  
 CC and metabolic. Polypeptides and polynucleotides of the invention are  
 CC useful in the treatment, or as a vaccine in the prevention of, cancer,  
 CC wound healing disorders, infection, atherosclerosis, Parkinson's disease  
 CC and Alzheimer's disease, autoimmune disorder, haematopoietic disorder,  
 CC inflammation, neoplastic diseases, nervous system related disorders and  
 CC cardiovascular disorders, pancreatitis, respiratory disorder,  
 CC hyperproliferation, systemic autoimmune disease, hyper-immunity,  
 CC developmental abnormality, gastrointestinal ulceration, neuropathy,  
 CC haematological diseases, metabolic diseases, sperm dysfunction, thyroid  
 CC disorders e.g. hypothyroidism, brain damages, colitis, cone photo-  
 CC transduction deficiency, neurological diseases, stroke, angiogenesis,  
 CC ovulation disorders, diseases in the spinal cord, thyroid gland, heart,  
 CC

CC trachea, thymus, lymph node and muscular system, obesity, anorexia,  
 CC growth abnormalities, and alleviation of precocious puberty, the  
 CC sequences given in records AB086130-AB086184 represent novel human cDNA's  
 CC of the invention.  
 XX Sequence 3066 BP; 738 A; 881 C; 852 G; 595 T; 0 other;  
 SQ Query Match 93.2%; Score 152.8; DB 24; Length 3066;  
 Best Local Similarity 95.7%; Pred. No. 4e-41;  
 Matches 157; Conservative 0; Mismatches 7; Indels 0; Gaps 0;  
 QY 1 GCACCGGACATCAAGCGAGGAATATTTTCTACTTGAAGAAGATAGAATCATGATGACAT 60  
 Db 735 GCACCGGACCTCAAGTCAGCAACATTTTCTACTTGAAGAAGATAGAATCATGATGACAT 794  
 QY 61 CTGCAATTAACATTTTCAAGATTACAGATTTTGGCTTGGCGAGGAGATGCGACAGGACAC 120  
 Db 795 CTGCAATTAACATTTTCAAGATTACAGATTTTGGCTTGGCGAGGAGATGCGACAGGACAC 854  
 QY 121 CAAATGAGCAGCAGCAGGACCATATGCTGATGCGCCCGCAGAG 164  
 Db 855 CAAATGAGCAGCAGCAGGACCATATGCTGATGCGCCCGCAGAG 898  
 RESULT 4  
 ABN86357  
 ID ABN86357 standard; DNA; 3111 BP.  
 XX AC ABN86357;  
 XX DT 08-OCT-2002 (first entry)  
 XX DE Novel human protein (NHP) kinase coding sequence.  
 XX KW Novel human protein; NHP; kinase; human; gene; ds.  
 XX OS Homo sapiens.  
 XX FH Key Location/Qualifiers  
 FT 1..3111  
 CDS /\*tag= a  
 FT /transl\_except= (pos: 2773..2775, aa: Xaa)  
 FT /product= "NHP kinase"  
 FT /note= "Xaa can be any amino acid"  
 FT  
 XX WO200255685-A2.  
 PN 18-JUL-2002.  
 PD  
 XX 10-DEC-2001; 2001WO-US47606.  
 PF 11-DEC-2000; 2000US-254744P.  
 PR (LEXI-) LEXICON GENETICS INC.  
 XX Hu Y, Kieke JA, Donoho G;  
 PI WPI; 2002-566739/60.  
 DR P-PSDB; ABB80923.  
 XX Novel human kinase polynucleotide encoding a protein that shares  
 PT structural similarity with animal kinases for therapeutic, diagnostic  
 PT and pharmacogenomic applications -  
 XX Claim 1; Page 36-37; 41pp; English.  
 XX The invention relates to a novel human protein (NHP), kinase that shares  
 CC structural similarity with animal kinases. The kinase polynucleotides are  
 CC useful in therapeutic, diagnostic and pharmacogenomic applications and  
 CC for identifying compounds that modulate, i.e. act as agonists or  
 CC antagonists of the gene expression or gene product activity. The present  
 CC sequence represents the NHP kinase coding sequence.  
 XX



used in the prevention, diagnosis and treatment of diseases cancers,  
 CC adenocarcinoma, leukaemia, sarcoma, immune disorder, Addison's disease,  
 CC acquired immune deficiency syndrome (AIDS), anaemia, asthma, allergies,  
 CC gout, microbial infections, cardiovascular disease and/or inflammation,  
 CC myasthenia gravis, atherosclerosis, cirrhosis, osteoporosis, myocardial  
 CC infarction, cataract, growth and development disorder, seizure disorder,  
 CC pulmonary embolism, Gaucher's disease, lipid disorder, lipid storage  
 CC disease, Pick's disease, Tay-Sachs disease, renal disease and obesity.  
 CC PKIN may be used to treat disorders associated with decreased PKIN  
 CC expression by rectifying mutations or deletions in a patient's genome  
 CC that affect the activity of PKIN by expressing inactive proteins or to  
 CC supplement the patient's own production of PKIN. PKIN nucleic acids may be  
 CC used to produce the PKIN polypeptide, by inserting the nucleic acids into  
 CC a host cell and culturing the cell to express the protein. PKIN nucleic  
 CC acid and its complementary sequences may also be used as DNA probes in  
 CC diagnostic assays to detect and quantitate the presence of similar  
 CC nucleic acid sequences in samples and therefore which patients may be  
 CC in need of restorative therapy. The present sequence is human PKIN-9  
 CC cDNA.

XX SQ Sequence 3141 BP; 685 A; 941 C; 942 G; 573 T; 0 other;

Query Match 57.4%; Score 94.2; DB 22; Length 3141;  
 Best Local Similarity 73.6%; Pred. No. 2.3e-21;  
 Matches 120; Conservative 0; Mismatches 43; Indels 0; Gaps 0;

QY 2 CACCGGACATCAAGCAGGAAATATTGCTACTTGAAGATAGACATGATGACATC 61  
 DB 766 CACCGGACCTTAACTCCAGCAACATATTGATCTCCAGAGGTTGGAGATGGAGCCTG 925  
 QY 62 TGCATTAACACTTTGAAGATTACAGATTTGGTTGGCGAGGGAATGGCAGACGACCC 121  
 DB 826 AGCAACAAGATTCTGAAGATCAGTATTTGGCTGGCTGGGAATGGCAGCAGACCC 885  
 QY 122 AAATGAGCAGCAGCAGGACCATGCTGCTGGATGCCCCCAGAG 164  
 DB 886 AAGATGAGTGGCGGAGGACGTATGCTTGGATGGCAGCCGGAAG 928

RESULT 7  
 AAD34309  
 ID AAD34309 standard; cDNA: 3538 BP.

XX AC AAD34309;

XX DT 16-JUL-2002 (first entry)

XX DE Human PKIN-12 cDNA.

XX KW Human; kinase; enzyme; PKIN-12 protein; immune system disorder; anaemia;  
 KW acquired immune deficiency syndrome; thymic hypoplasia; Crohn's disease;  
 KW asthma; neurological disorder; epilepsy; Charcot-Marie-Tooth disease;  
 KW AIDS; seizures; cell proliferative disorder; cancer; adenocarcinoma;  
 KW leukaemia; lymphoma; melanoma; myeloma; sarcoma; developmental disorder;  
 KW Down's syndrome; gene therapy; protein therapy; cytostatic; gene; ss.

XX OS Homo sapiens.

XX FH Key Location/Qualifiers

FT CDS 1..3294

FT /tag= a

FT /product= "Human PKIN-12 protein"

FT sig\_peptide 1..51

FT /tag= b

FT mat\_peptide 52..3291

FT /tag= c

FT /product= "Human mature PKIN-12 protein"

XX PN W0200218557-A2.

XX PD 07-MAR-2002.

XX PF 31-AUG-2001; 2001WO-US27219.

XX PR 31-AUG-2000; 2000US-229873P.  
 PR 08-SEP-2000; 2000US-231357P.  
 PR 14-SEP-2000; 2000US-232654P.  
 PR 22-SEP-2000; 2000US-234902P.  
 PR 29-SEP-2000; 2000US-236499P.  
 PR 06-OCT-2000; 2000US-238389P.  
 PR 13-OCT-2000; 2000US-240542P.

XX (INCY-) INCYTE GENOMICS INC.

XX PA Bandman O, Nguyen DB, Wallia NK, Hafalia AJA, Yao MG, Gandhi AR;

XX PI Gururajan R, Ding L, Patterson C, Yue H, Baughn MR, Tribouley CM;

XX PI Thornton M, Elliott VS, Lu Y, Ison CH, Au-Young J, Tang Y;

XX PI Azimzai Y, Burilli JD, Marcus GA, Zingler KA, Lu DM, Lal PG;

XX PI Ramkumar J, Warren BA, Kearney L, Pollicky JL, Thangavelu K;

XX PI Burford N;

XX WPI; 2002-329769/36.

XX P-PSDB; AAE21717.

XX New human kinases, useful for diagnosing, treating or preventing immune

XX system disorders (e.g. Crohn's disease), neurological disorders (e.g.

XX epilepsy), or cell proliferative disorders (e.g. cancers such as

XX leukemia or lymphoma)

XX Claim 91; Page 207-208; 218pp; English.

XX The present invention relates to human kinases (PKIN) and polynucleotides

XX encoding such proteins. PKIN sequences of the invention are useful for

XX diagnosing, treating or preventing disorders associated with aberrant

XX expression of PKIN, particularly immune system disorders (e.g. acquired

XX immune deficiency syndrome (AIDS), thymic hypoplasia, Crohn's disease,

XX anaemia, asthma), neurological disorders (e.g. epilepsy, Charcot-Marie-

XX Tooth disease or seizures), cell proliferative disorders (e.g. cancers

XX such as adenocarcinoma, leukaemia, lymphoma, melanoma, myeloma, sarcoma),

XX and developmental disorders (e.g. Down's syndrome). They are also used

XX in gene therapy and protein therapy. The present sequence is a cDNA

XX encoding human PKIN-12 protein.

XX SQ Sequence 3538 BP; 763 A; 1055 C; 1062 G; 658 T; 0 other;

Query Match 57.4%; Score 94.2; DB 24; Length 3538;

Best Local Similarity 73.6%; Pred. No. 2.4e-21;

Matches 120; Conservative 0; Mismatches 43; Indels 0; Gaps 0;

QY 2 CACCGGACATCAAGCAGGAAATATTGCTACTTGAAGATAGACATGATGACATC 61

DB 796 CACCGGACCTTAACTCCAGCAACATATTGATCTCCAGAGGTTGGAGATGGAGCCTG 855

QY 62 TGCATTAACACTTTGAAGATTACAGATTTGGTTGGCGAGGGAATGGCAGACGACCC 121

DB 856 AGCAACAAGATTCTGAAGATCAGTATTTGGCTGGCTGGGAATGGCAGCAGACCC 915

QY 122 AAATGAGCAGCAGCAGGACCATGCTGCTGGATGCCCCCAGAG 164

DB 916 AAGATGAGTGGCGGAGGACGTATGCTTGGATGGCAGCCGGAAG 958

RESULT 8

AAI85458/C

ID AAI85458 standard; cDNA: 444 BP.

XX AC AAI85458;

XX DT 06-NOV-2001 (first entry)

XX DE Human polynucleotide SEQ ID NO 5518.

XX KW Human; cytokine; cell proliferation; cell differentiation; gene therapy;

XX vaccine; peptide therapy; stem cell growth factor; haematopoiesis;

XX tissue growth factor; immunomodulatory; cancer; leukaemia;

XX nervous system disorders; arthritis; inflammation; ss.



|    |   |  |  |
|----|---|--|--|
| XX | Seq   | Sequence 3558 BP; 635 A; 1211 C; 1161 G; 551 T; 0 other;           |  |
|    | Query Match   | 50.6%; Score 93; DB 24; Length 3558;                               |  |
|    | Best Local Similarity   | 69.3%; Pred. No. 1.4e-17;  |  |
|    | Matches 113; Conservative   | 0; Mismatches 50; Indels 0; Gaps                                   |  |
| QY | 2   | CACCGGACATCAGGAGGAGAAATATTTTGGCTACTTGGAGAGATAGAACATGATGACATC 61    |  |
| DB | 1196  | CACCGTGATCTCAAGTCCACACAACTTTTGGCTGTCGAGCCCATTTGAGAGTGAGGACATG 1255 |  |
| QY | 62  | TGCAATAAAACCTTTGAAGATTACAGATTTTGGTTGGCGAGGGAATGGCAGACAGACC 121     |  |
| DB | 1296  | GAGCACAGACCCCTGAAGATCACCGGACTTTGGCTGGCCGAGTGCGCAGAAACACCA 1315     |  |
| QY | 122   | AAATGAGCACAGGACGACCTATGCTGGATGGCCCGCAGAG 164                       |  |
| DB | 1316  | CAAATGAGTGGCGGGCAGCTACGCTCGATGCTCTCCTGAGG 1358                     |  |
|    | RESULT 11   |  |  |
| ID | ABL70018  |  |  |
| AC | ABL70018 standard; DNA: 3454 BP.  |  |  |
| AC | ABL70018;   |  |  |
| DT | 15-MAY-2002 (first entry)   |  |  |
| DE | Pancreas cancer related gene sequence SEQ ID NO:8355.                   |  |  |
| KW | Human; cancer; colon; breast; ovary; oesophagus; kidney; thyroid;       |  |  |
| KW | Stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous;    |  |  |
| KW | Cystosatic; gene therapy; antineoplastic; Wilms tumour; adenocarcinoma; |  |  |
| OS | gene; ds.   |  |  |
| OS | Homo sapiens.   |  |  |
| PN | WO200194629-A2.   |  |  |
| PD | 13-DEC-2001.  |  |  |
| PF | 30-MAY-2001; 2001WO-US10838.  |  |  |
| XX | 05-JUN-2000; 2000US-209473P.  |  |  |
| PR | 05-JUN-2000; 2000US-209531P.  |  |  |
| PR | 18-SEP-2000; 2000US-233133P.  |  |  |
| PR | 18-SEP-2000; 2000US-233617P.  |  |  |
| PR | 20-SEP-2000; 2000US-234009P.  |  |  |
| PR | 20-SEP-2000; 2000US-234034P.  |  |  |
| PR | 20-SEP-2000; 2000US-234052P.  |  |  |
| PR | 22-SEP-2000; 2000US-234509P.  |  |  |
| PR | 22-SEP-2000; 2000US-234567P.  |  |  |
| PR | 25-SEP-2000; 2000US-234923P.  |  |  |
| PR | 25-SEP-2000; 2000US-234934P.  |  |  |
| PR | 25-SEP-2000; 2000US-235077P.  |  |  |
| PR | 25-SEP-2000; 2000US-235082P.  |  |  |
| PR | 25-SEP-2000; 2000US-235134P.  |  |  |
| PR | 25-SEP-2000; 2000US-235280P.  |  |  |
| PR | 26-SEP-2000; 2000US-235637P.  |  |  |
| PR | 26-SEP-2000; 2000US-235638P.  |  |  |
| PR | 27-SEP-2000; 2000US-235711P.  |  |  |
| PR | 27-SEP-2000; 2000US-235720P.  |  |  |
| PR | 27-SEP-2000; 2000US-235840P.  |  |  |
| PR | 28-SEP-2000; 2000US-235863P.  |  |  |
| PR | 28-SEP-2000; 2000US-236028P.  |  |  |
| PR | 28-SEP-2000; 2000US-236032P.  |  |  |
| PR | 28-SEP-2000; 2000US-236033P.  |  |  |
| PR | 28-SEP-2000; 2000US-236034P.  |  |  |
| PR | 28-SEP-2000; 2000US-236109P.  |  |  |
| PR | 28-SEP-2000; 2000US-236111P.  |  |  |
| PR | 29-SEP-2000; 2000US-236842P.  |  |  |
| PR | 29-SEP-2000; 2000US-236891P.  |  |  |
| PR | 02-OCT-2000; 2000US-237172P.  |  |  |

PR 02-OCT-2000; 2000US-237173P.  
PR 02-OCT-2000; 2000US-237278P.  
PR 02-OCT-2000; 2000US-237295P.  
PR 02-OCT-2000; 2000US-237295P.  
PR 02-OCT-2000; 2000US-237316P.  
PR 03-OCT-2000; 2000US-237425P.  
PR 03-OCT-2000; 2000US-237598P.  
PR 03-OCT-2000; 2000US-237604P.  
PR 03-OCT-2000; 2000US-237606P.  
PR 03-OCT-2000; 2000US-237608P.  
PR 01-NOV-2000; 2000US-244867P.  
PR 01-NOV-2000; 2000US-245084P.  
XX (AVAL-) AVALON PHARM.  
XX PA  
XX PI Young PE, Augustus M, Carter KC, Ebner R, Endress G, Horrigan S;  
XX PI Soppet DR, Weaver Z;  
XX XX  
XX DR WPI; 2002-188264/24.  
XX XX  
XX PT Screening for anti-neoplastic agent involves exposing cells to a  
XX PT chemical agent to be tested for anti-neoplastic activity, and  
XX PT determining a change in expression of a gene of a signature gene set -  
XX XX  
XX PS Claim 1; SEQ ID 8355; 44pp; English.  
XX XX  
XX CC The present invention describes a method (M1) for screening for an  
XX CC anti-neoplastic agent. The method involves exposing cells to a chemical  
XX CC agent to be tested for anti-neoplastic activity, determining a change in  
XX CC expression of at least one gene (1) of a signature gene set, where (1)  
XX CC comprises a sequence (S) selected from 847 sequences (given in AB51564  
XX CC to AB170110) or is at least 95% identical to (S) where a change in  
XX CC expression is indicative of anti-neoplastic activity. (1) has cytostatic  
XX CC activity and can be used in gene therapy. M1 can be used for screening  
XX CC an anti-neoplastic agent, and can be used for producing a product which  
XX CC is the data collected with respect to the anti-neoplastic agent as a  
XX CC result of M1 and the data is sufficient to convey the chemical  
XX CC structure and/or properties of the agent. M1 can be used in the  
XX CC treatment of cancer such as colon, breast, stomach, lung, thyroid,  
XX CC oesophageal, ovarian, kidney, prostate or pancreatic cancer,  
XX CC adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer,  
XX CC infiltrating lobular cancer, squamous cell carcinoma, neuroendocrine  
XX CC carcinoma, papillary carcinoma and Wilms' tumour.  
XX SQ Sequence 3454 BP; 594 A; 1217 C; 1136 G; 507 T; 0 other;  
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Best Local Similarity 65.68; Pred. No. 2.5e-14;  
Matches 107; Conservative 0; Mismatches 56; Indels 0; Gaps 0;  
QY 2 CACGGGACATCAGCGAGGAGATATTTGCTACTTGAGAGATAGACATGATCATC 61  
DB 946 CACGGGAGCTTCACTCATCATCTCTGATCTGAGGAGGAGGAGGAGGAGGAGG 1005  
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DB 1006 GCAGACCGGTCTCAGATGACGAGCTTCGGCCCTCGCCGCGGAGGAGGAGGAGGAGG 1065  
QY 122 AATATGACAGACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 164  
DB 1066 AAGATGAGGCTCGCGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 1108  
RESULT 12  
ABAL177  
ID ABAL177 standard; DNA: 7084 BP.  
XX AC  
XX AC ABAL177;  
XX XX  
XX DT 23-JAN-2002 (first entry)  
XX DT Human nervous system related polynucleotide SEQ ID NO 9508.  
XX DE  
XX XX

KW Human; nootropic; neuroprotective; cytostatic; dermatological; virucide;  
KW immunosuppressive; anti-inflammatory; anti-HIV; antibacterial; vulnerary;  
KW antiparkinsonian; antiskilling; antianaemic; antiarthritic; cancer;  
KW antirheumatic; hepatotropic; cerebroprotective; antiinflammatory;  
KW antiallergic; antidiabetic; antitumor; anticonvulsant; antifungal;  
KW antiparasitic; cardiast; immune disorder; cardiovascular disorder;  
KW neurological disease; infection; nephrotropic; gene therapy; vaccine; ds.  
XX Homo sapiens.  
XX OS  
XX PN WO200159063-A2.  
XX XX  
XX PD 16-AUG-2001.  
XX XX  
XX PF 17-JAN-2001; 2001WO-US01334.  
XX XX  
XX PR 31-JAN-2000; 2000US-0179065.  
XX PR 04-FEB-2000; 2000US-0180628.  
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XX PR 14-SEP-2000; 2000US-0232400.  
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XX PD 09-AUG-2001.  
XX XX 17-JAN-2001; 2001WO-US01354.  
PF XX  
XX  
PR 31-JAN-2000; 2000US-0179065.  
PR 04-FEB-2000; 2000US-0180628.  
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PR 01-DEC-2000; 2000US-0250160.  
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PR 08-DEC-2000; 2000US-0251989.  
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PR 11-DEC-2000; 2000US-0254097.  
PR 05-JAN-2001; 2001US-02559678.  
XX  
XX (HUMA-) HUMAN GENOME SCI INC.  
XX PA  
XX PI Rosen CA, Barash SC, Ruben SM;  
XX XX

DR WPI; 2001-483426/52.  
 XX Nucleic acids encoding human immune/hematopoietic antigen polypeptides,  
 PT useful for preventing, diagnosing and/or treating cancers and  
 PT metastasis.  
 XX  
 PS Disclosure; SEQ ID NO 28728; 3071pp + Sequence Listing; English.  
 XX  
 CC AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I)  
 CC amino acid sequences given in AAM82170 to AAM9121. (I) have cytostatic  
 CC activity, and can be used in gene therapy and vaccine production. (I)  
 CC proteins and polynucleotides may be used in the prevention, diagnosis and  
 CC treatment of diseases associated with inappropriate (I) expression. For  
 CC example, they may be used to treat disorders associated with decreased  
 CC expression by rectifying mutations or deletions in a patient's genome  
 CC that affect the activity of (I) by expressing inactive proteins or to  
 CC supplement the patient's own production of (I). Additionally, (I)  
 CC polynucleotides may be used to produce the secreted (I), by inserting  
 CC the nucleic acids into a host cell and culturing the cell to express the  
 CC protein. (I) proteins and polynucleotides may be used to prevent,  
 CC diagnose and treat immune/haematopoietic-related diseases, especially  
 CC cancers and cancer metastases of haematopoietic-derived cells. AAK64703  
 CC to AAK87694 represent human immune/haematopoietic antigen genomic  
 CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169  
 CC represent sequences used in the exemplification of the present invention.  
 XX  
 SQ Sequence 7084 BP; 1519 A; 1987 C; 2073 G; 1505 T; 0 other;  
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 Matches 97; Conservative 0; Mismatches 41; Indels 0; Gaps 0;  
 QY 27 TTTTGTCTACTGAGAGATGACATGACATCTGCAATAAATTTGAAGATTACAG 86  
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 QY 87 ATTTTGGTGTGGCAGGAGATGACAGCAGGACCAACCAATGAGCAGCAGGACCTATG 146  
 DB 4730 ACTTTGGCTGTGGCCGAGAGTGACCAAAACCAACCAATGAGTGCCTGGCGGACCTACG 4789  
 QY 147 CCTGGATGGCCCGCAGAG 164  
 DB 4790 CCTGGATGGCTCTGTAGG 4807  
 RESULT 14  
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 AC AAK49755;  
 XX  
 DT 10-MAR-1994 (first entry)  
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 DE pTK gene LpTK4 partial sequence.  
 XX  
 KW pTK; protein tyrosine kinase; catalytic domain; c-kit; megakaryocyte;  
 KW lymphocyte; amplification; primer; polymerase chain reaction; PCR; ds.  
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PD 05-AUG-1993.  
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 PF 22-JAN-1993; 93WO-US00586.  
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 PR 22-JAN-1992; 92US-0826935.  
 XX  
 PA (NEW-) NEW ENGLAND DEACONESS HOSPITAL.  
 XX  
 PI Avraham H, Cowley S, Groopman J, Scadden D;  
 XX WPI; 1993-320330/40.  
 XX  
 DR New protein tyrosine kinase genes and proteins encoded by genes -  
 PT are of human mega-karyocytic origin  
 XX  
 PS Claim 2; Fig 6; 60pp; English.  
 XX  
 CC pTK genes were identified using two sets of degenerative  
 CC oligonucleotide primers: a first set which amplifies all pTK DNA  
 CC segments (AAQ49743-44), and a second set which amplifies highly  
 CC conserved sequences present in the catalytic domain of the c-kit  
 CC subgroup of pTKs (AAQ49745-46). The pTK genes identified are described  
 CC in AAQ49747-57 and AAK41897-02.  
 CC The LpTKs are expressed in lymphocytic cells, as well as  
 CC megakaryocytic cells. Sequencing of LpTK-4 revealed the sequences  
 CC given in AAQ49751 and AAQ49755. The protein sequence corresp. to  
 CC AAQ49751 is claimed (claim 7) and stated as given in the specification,  
 CC however is missing from the publication.  
 XX  
 SQ Sequence 404 BP; 95 A; 106 C; 123 G; 77 T; 3 other;  
 Query Match 43.9%; Score 72; DB 14; Length 404;  
 Best Local Similarity 59.6%; Pred. No. 2.9e-14;  
 Matches 96; Conservative 0; Mismatches 42; Indels 0; Gaps 0;  
 QY 27 TTTTGTCTACTGAGAGATGACATGACATCTGCAATAAATTTGAAGATTACAG 86  
 DB 103 TTTTGTCTGTCAGCCATTTGAGAGTGACGACATGAGCACAAGACCTGAAGATCACC 162  
 QY 87 ATTTTGGTGTGGCAGGAGATGACAGCAGGACCAACCAATGAGCAGCAGGACCTATG 146  
 DB 163 ACTTTGGCTGTGGCCGAGAGTGACCAAAACCAACCAATGAGTGCCTGGCGGACCTACN 222  
 QY 147 CCTGGATGGCCCGCAGAG 164  
 DB 223 CCTGGATGGCTCTGTAGG 240  
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 ID AAT03098 standard; DNA; 404 BP.  
 AC AAT03098;  
 XX  
 DT 14-FEB-1996 (first entry)  
 XX  
 DE Protein tyrosine-kinase LpTK4 DNA fragment.  
 XX  
 KW Protein tyrosine-kinase; pTK; LpTK4; agonist; cell growth;  
 KW differentiation; ss.  
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 OS Homo sapiens.  
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XX      12-OCT-1995.
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XX      04-APR-1994; 94US-0222616.
XX
XX      (GETH ) GENENTECH INC.
XX
XX      Bennett BD, Goeddel D, Lee JM, Matthews W, Tsai SP;
PI      Wood WI;
XX
XX      WPI; 1995-366160/47.
XX
XX      Agonist antibodies which activate specific protein tyrosine
XX      kinase(s) - also activate chimeric proteins of kinase extracellular
XX      domain and Ig constant domain, useful for studying, and therapeutic
XX      modulation of, cell growth and differentiation
XX
XX      Disclosure; Page 58-59; 125pp; English.
XX
XX      DNA probes based on protein tyrosine-kinase (ptk) sequences were used
XX      to screen cDNA libraries to identify novel ptk genes. A LptK4 gene
XX      partial sequence (AAT03098) was isolated from a lymphocytic cell
XX      library. The partial sequence can be used to identify other new
XX      ptk genes, or to design drugs, peptides or antisense constructs
XX      that modulate ptk activity.
XX
XX      Sequence 404 BP; 95 A; 106 C; 123 G; 77 T; 3 other;

Query Match      43.9%; Score 72; DB 16; Length 404;
Best Local Similarity 69.6%; Pred. No. 2.9e-14;
Matches 96; Conservative 0; Mismatches 42; Indels 0; Gaps 0;

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DB      ||||| || || || || || || || || || || || || || || || ||
DB      103 TTTTCTCTGTCACCCCATTTGAGAGTGAGCAGATGAGCAGACCAAGACCTGAAGATCACCG 162
QY      87 ATTTGGGTTGGCGAGGGAATGGCAGGAGGACCAACCAATGAGCAGCAGGCGACCTATG 146
DB      ||||| ||||| || || || || || || || || || || || || || || || ||
DB      163 ACTTTGGCCTGGCCCGAGAGTGGCACAACCAACCAACACACAAATGAGTGCCTCCNGSCACCTACN 222
QY      147 CCTGATGGCCCCCAGAG 164
DB      ||||| ||||| || || ||
DB      223 CCTGGATGGCTCCTGAGG 240
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